## II. CLAIMS

1. (Currently Amended) A  $\frac{G_{\alpha q \; Gustducin}}{G16/gust \; 44}$  or  $\frac{G15/gust44}{G16/gust \; 44}$  or  $\frac{G16/gust \; 44}{G16/gust \; 44}$  or  $\frac{G15/gust44}{G16/gust \; 44}$  protein sequence are replaced with a 44 amino acid unit of Gustducin, where such 44 amino acid unit of Gustducin is the last 44 amino acids of SEQ ID NO:2.

## 2-5. Cancelled

- 6. (Previously Presented) A G-protein according to claim 1 encoded for by the nucleic acid set forth in SEQ ID NO:1.
- 7. (Previously Presented) A nucleic acid comprising the nucleotide sequence set forth in SEQ ID NO:1 encoding for a G-protein according to claim 1.
- 8. (Previously Presented) An expression vector comprising nucleic acid comprising the nucleotide sequence set forth in SEQ ID NO:1 encoding for a G-protein according to claim 1.
- 9. (Previously Presented) A host cell transformed with an expression vector according to claim 8.
- 10. (Previously Presented) A method of producing a chimeric Gprotein according to claim 1 comprising the step of culturing
  host cells having contained therein an expression vector
  encoding for the chimeric G-protein, under conditions sufficient
  for expression of said G-protein, thereby causing production of
  the protein, and recovering the protein produced by the cell.

- 11. (Previously Presented) A method of analysis and discovery of modulators of bitter taste receptors using the chimeric proteins according to defined in claim 1.
- 12. (Previously Presented) A method according to claim 11 employing a mammalian cell-based assay employing a transfected gene or cDNA encoding a chimeric protein of the invention and a taste receptor, the method comprising the steps of contacting a compound with cells, and determining the functional effect of the compound on chimeric G-protein.
- 13. Previously Presented) A method according to claim 10 wherein the functional effect is determined by measuring the changes in intracellular messengers IP3 or calcium<sup>2+</sup>.

## 14-17. Cancelled

- 18. (Currently Amended) A  $G_{\alpha q \; Gust \; ducin}$  G16/gust 44 or G15/gust44 chimeric G-protein wherein the last 44 amino acids of the  $G_{\alpha q}$  G16/gust 44 or G15/gust44 protein sequence are replaced with a 44 amino acid unit of Gustducin, where such 44 amino acid unit of Gustducin is the last 44 amino acids of SEQ ID NO:2, and wherein the resulting  $G_{\alpha q gust \; 44}$  chimeric G-protein has a sequence homology of at least 80% in the last 44 amino acids of SEQ ID NO:2.
- 19. (Previously Presented) The chimeric G-protein of claim 18 having a sequence homology of at least 90% in the last 44 amino acids of SEQ ID NO:2.
- 20. (Previously Presented) The chimeric G-protein of claim 18

having a sequence homology of at least 95% in the last 44 amino acids of SEQ ID NO:2.

- 21. (Currently Amended) A  $\frac{G_{\alpha q \; Custducin}}{G16/gust \; 44}$  or  $\frac{G15/gust44}{gust \; 6-protein}$  wherein the last 44 amino acids of the  $\frac{G_{\alpha q}}{G16/gust \; 44}$  or  $\frac{G15/gust44}{gust \; 6-gust6}$  protein sequence are replaced with a 44 amino acid unit of Gustducin, where such 44 amino acid unit of Gustducin is the last 44 amino acids of SEQ ID NO:2, and wherein the resulting  $G_{\alpha q-gust44}$  chimeric G-protein has a sequence homology of at least 80% to SEQ ID NO:2.
- 22. (Previously Presented) The chimeric G-protein of claim 21 having a sequence homology of at least 90% to SEQ ID NO:2.
- 23. (Previously Presented) The chimeric G-protein of claim 21 having a sequence homology of at least 95% to SEQ ID NO:2.
- 24. (Currently Amended) A  $G_{aq\ Gustducin}$  G16/gust 44 or G15/gust44 chimeric G-protein wherein the last 44 amino acids of the  $G_{aq}$  G16/gust 44 or G15/gust44 protein sequence are replaced with a 44 amino acid unit of Gustducin, where such 44 amino acid unit of Gustducin is the last 44 amino acids of SEQ ID NO:2, and wherein the resulting  $G_{aq\ gust44}$  chimeric G-protein has a sequence homology of at least 80% to SEQ ID NO:2 and the chimeric protein binds to one or more of the human bitter, sweet and umami taste receptors.
- 25. (Previously Presented) The chimeric G-protein of claim 24 having a sequence homology of at least 90% to SEQ ID NO:2.

26. (Previously Presented) The chimeric G-protein of claim 24 having a sequence homology of at least 95% to SEQ ID NO:2.

## 27. Canceled

- 28. (Previously Presented) A nucleic acid encoding for a G-protein according to claim 18.
- 29. (Previously Presented) An expression vector comprising nucleic acid comprising the nucleotide sequence encoding for a G-protein according to claim 18.
- 30. (Previously Presented) A host cell transformed with an expression vector according to claim 29.
- 31. (Previously Presented) A method of producing a chimeric Gprotein according to claim 18 comprising the step of culturing
  host cells having contained therein an expression vector
  encoding for the chimeric G-protein, under conditions sufficient
  for expression of said G-protein, thereby causing production of
  the protein, and recovering the protein produced by the cell.
- 32. (Previously Presented) A method of analysis and discovery of modulators of bitter taste receptors using the chimeric proteins according to defined in claim 18.
- 33. (Previously Presented) A method according to claim 32 employing a mammalian cell-based assay employing a transfected gene or cDNA encoding a chimeric protein of the invention and a taste receptor, the method comprising the steps of contacting a compound with cells, and determining the functional effect of

USSN 10/538,038

Response to Office Action dated October 28, 2008

Atty. Docket: 102790-135

the compound on chimeric G-protein.

34. (Previously Presented)) A method according to claim 31 wherein the functional effect is determined by measuring the changes in intracellular messengers IP3 or calcium<sup>2+</sup>.